

Figure 1

1 RTPSDKPVAH VVANPQAEQG LQWLNRRANA LLANGVELRD NQLVVPSEGL
51 YLIYSQVLEK GQGCPSTHVL LTHTISRIV SYQTKVNLIS AIKSPCQRET
101 PEGAEAKPWY EPIYLGGVFQ LEKGDRLSAE INRPDYLLFA ESGQVYFGII
151 AL (SEQ ID NO: 6)

Figure 2

1 HSDPARRGQL SVCDSISEWV TAADKKTAVD MSGGTVTVLE KVPVSKGQLK
51 QYFYETKCNP MGYTKEGCRG IDKRHWNSQC RTTQSIYVRAL TMDSKKRIGW
101 RFIRIDTSCV CTLTIKRGR (SEQ ID NO: 7)

FIGURE 3

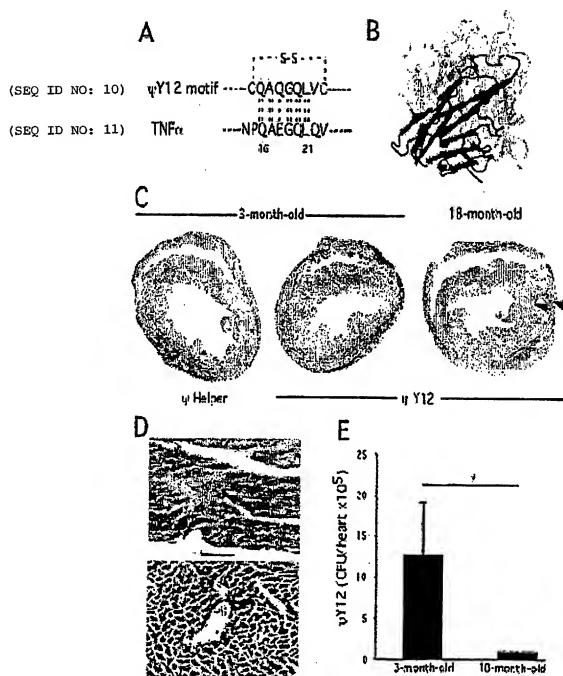


FIGURE 6

